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FIGURE 1

SK-N-MC  
SM1444  
KC1445  
U937

SK-N-MC  
SM1444  
SM1444  
KC1445

U937

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- 23

- 9.4

- 6.6

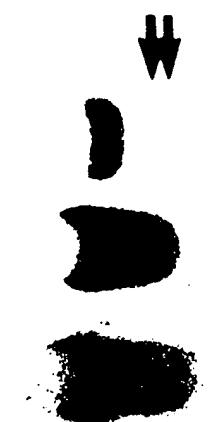


$C_{\mu}$

Hind III

$C_{\mu}$   $J_H$   $C_{\mu}$   $C_{\mu}$

BamH I



$C_{\mu}$   $J_H$   $C_{\mu}$   $C_{\mu}$

zyme

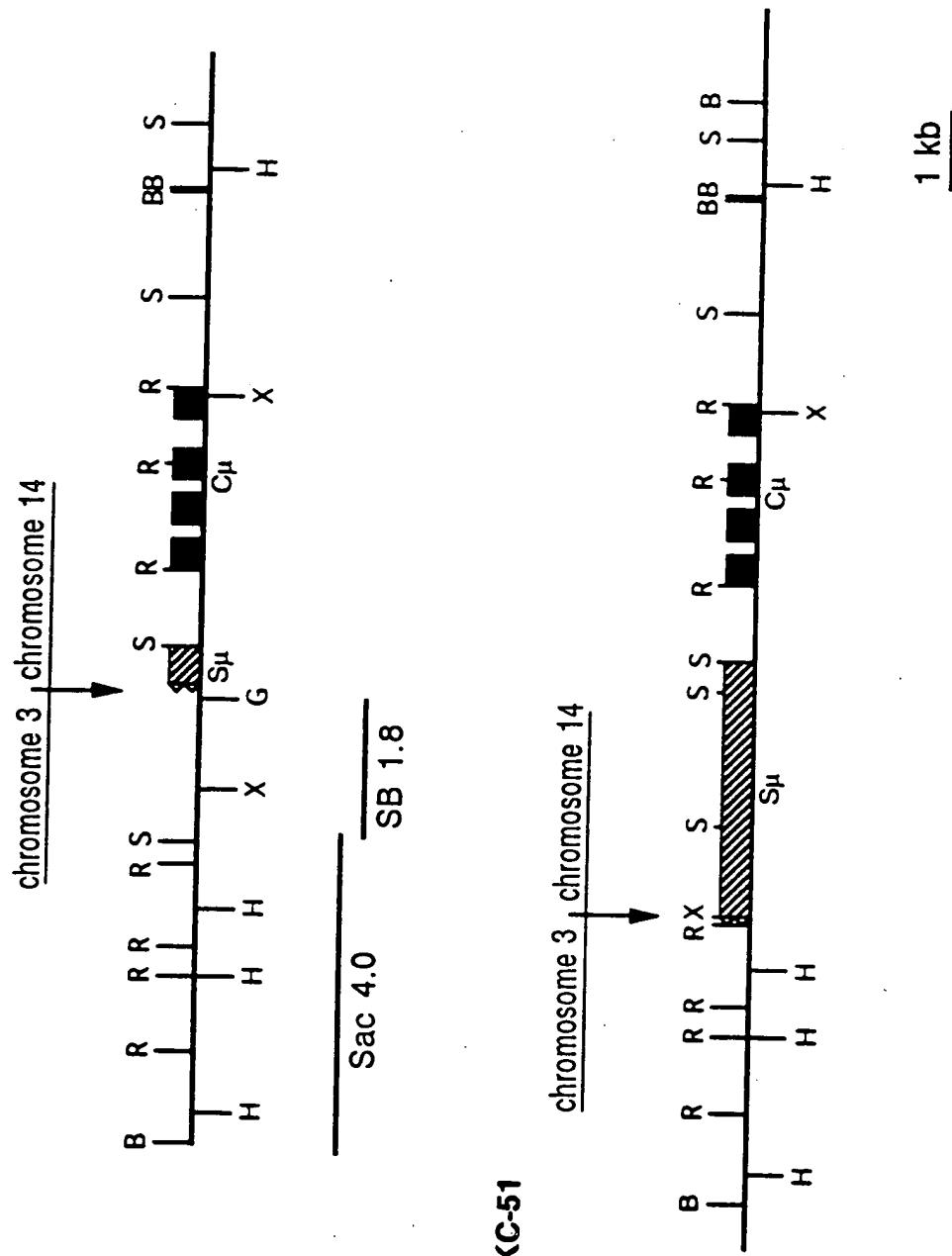


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**FIGURE 2** **SM-71**



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FIGURE 3B

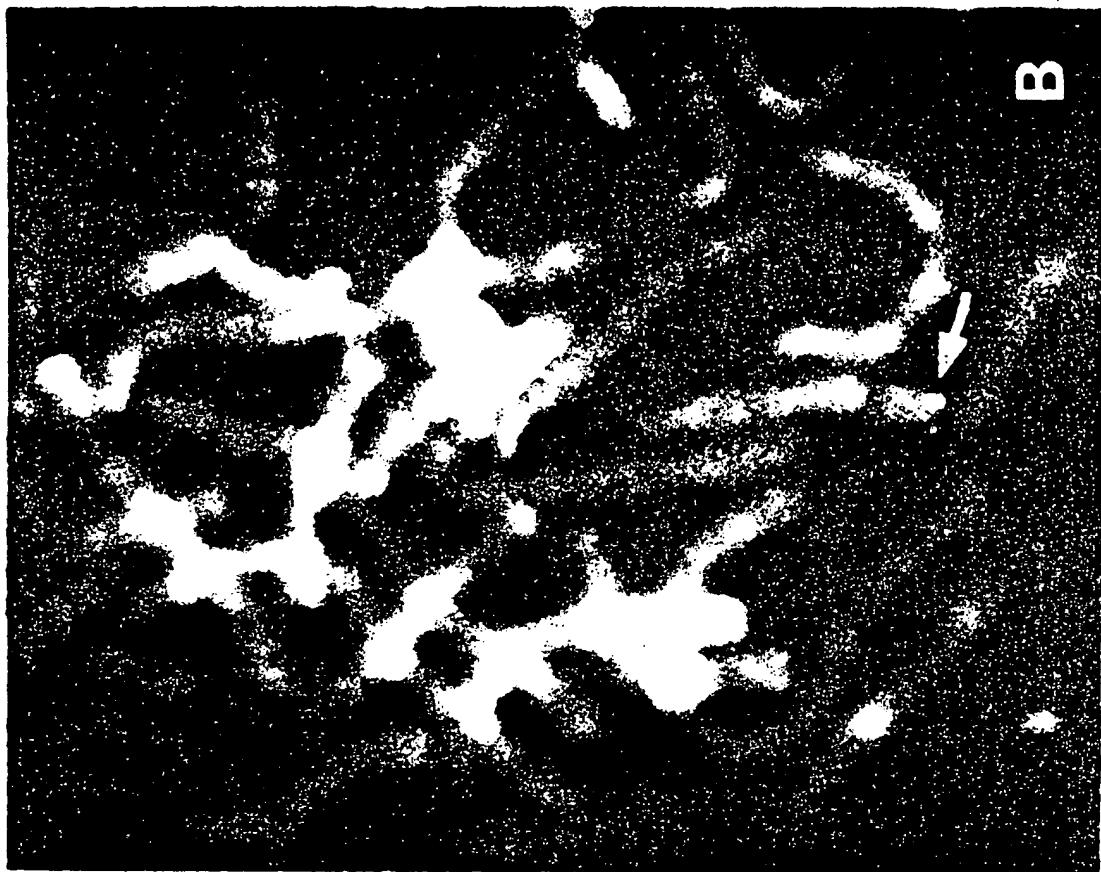


FIGURE 3A



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FIGURE 4A

FIGURE 4B

FIGURE 4C

SK-N-MC  
KC1445  
SM1444  
LD1411  
TF1403  
U937

23 —  
9.4 —  
6.6 —  
4.4 —

2.3 —

Enzyme  
probe  
*EcoR I*  
*SBI.8*

SK-N-MC

KC1445

SM1444

LD1411

TF1403

U937

23 —

9.4 —  
6.6 —  
4.4 —

2.3 —

*Bgl II*  
*Sac 4.0*

SK-N-MC

SO955

CF755

EM352

U937

23 —

9.4 —  
6.6 —  
4.4 —

2.3 —  
2.0 —

*Bgl II*  
*Sac 4.0*

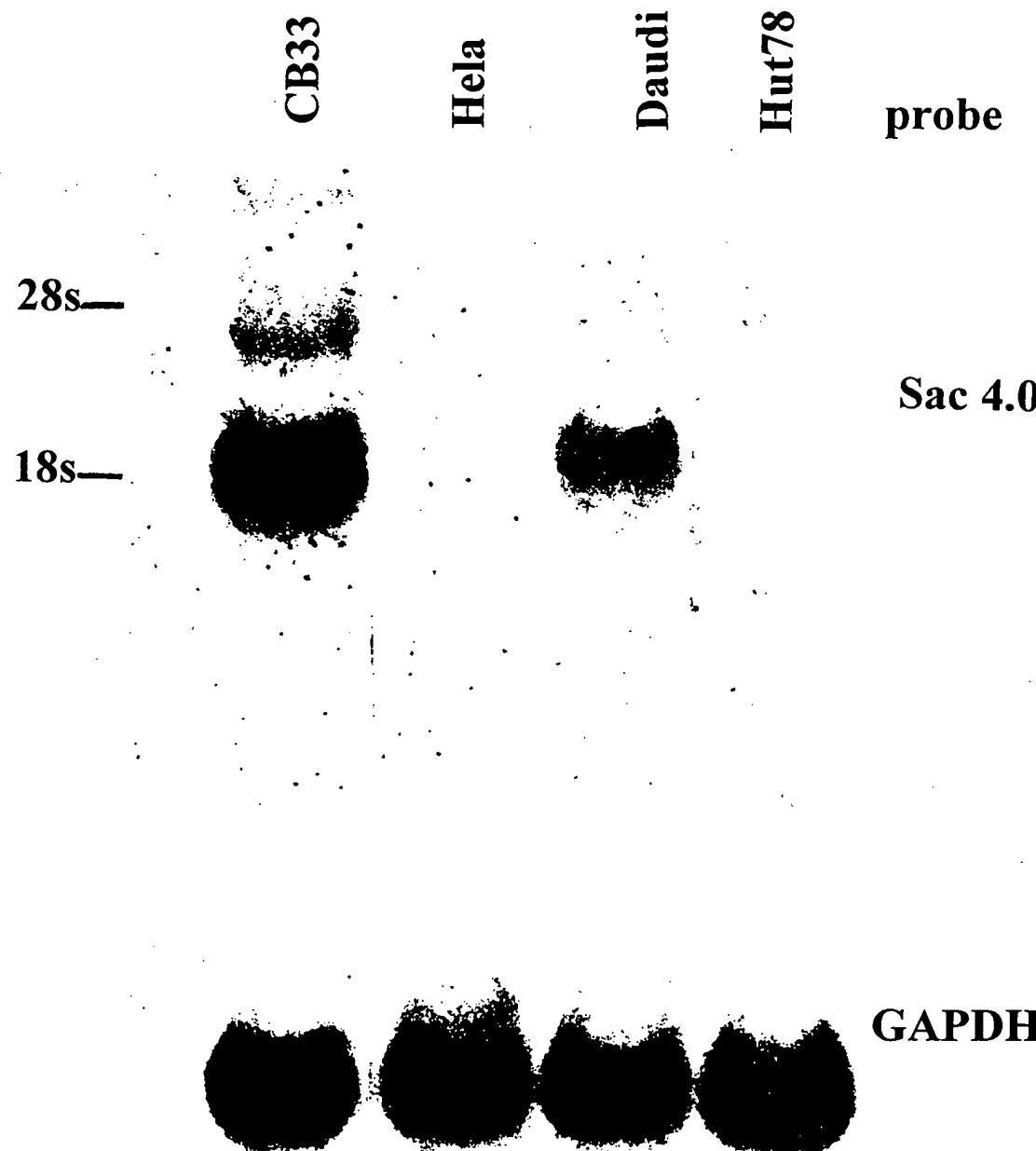
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**FIGURE 5**



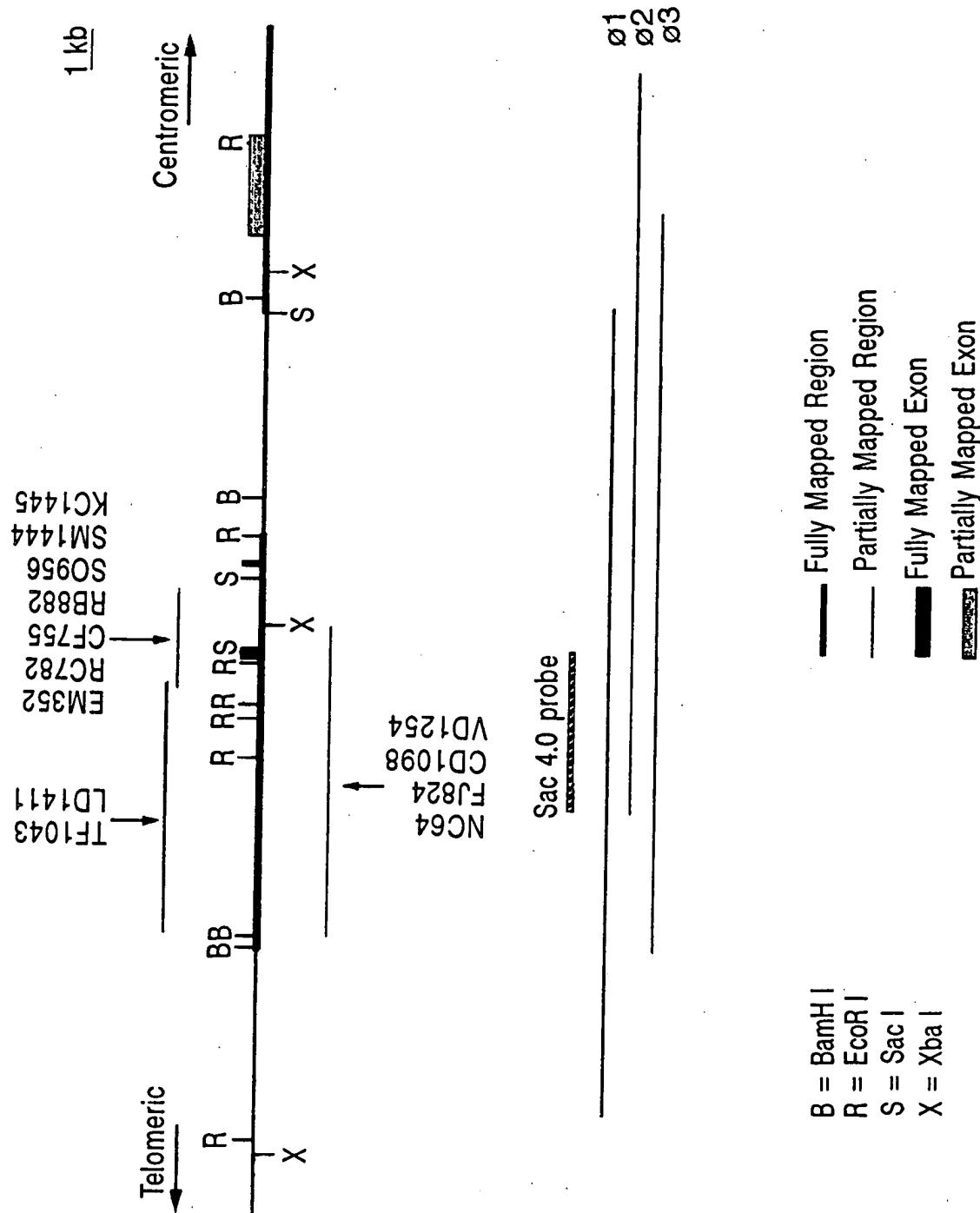


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FIGURE 6

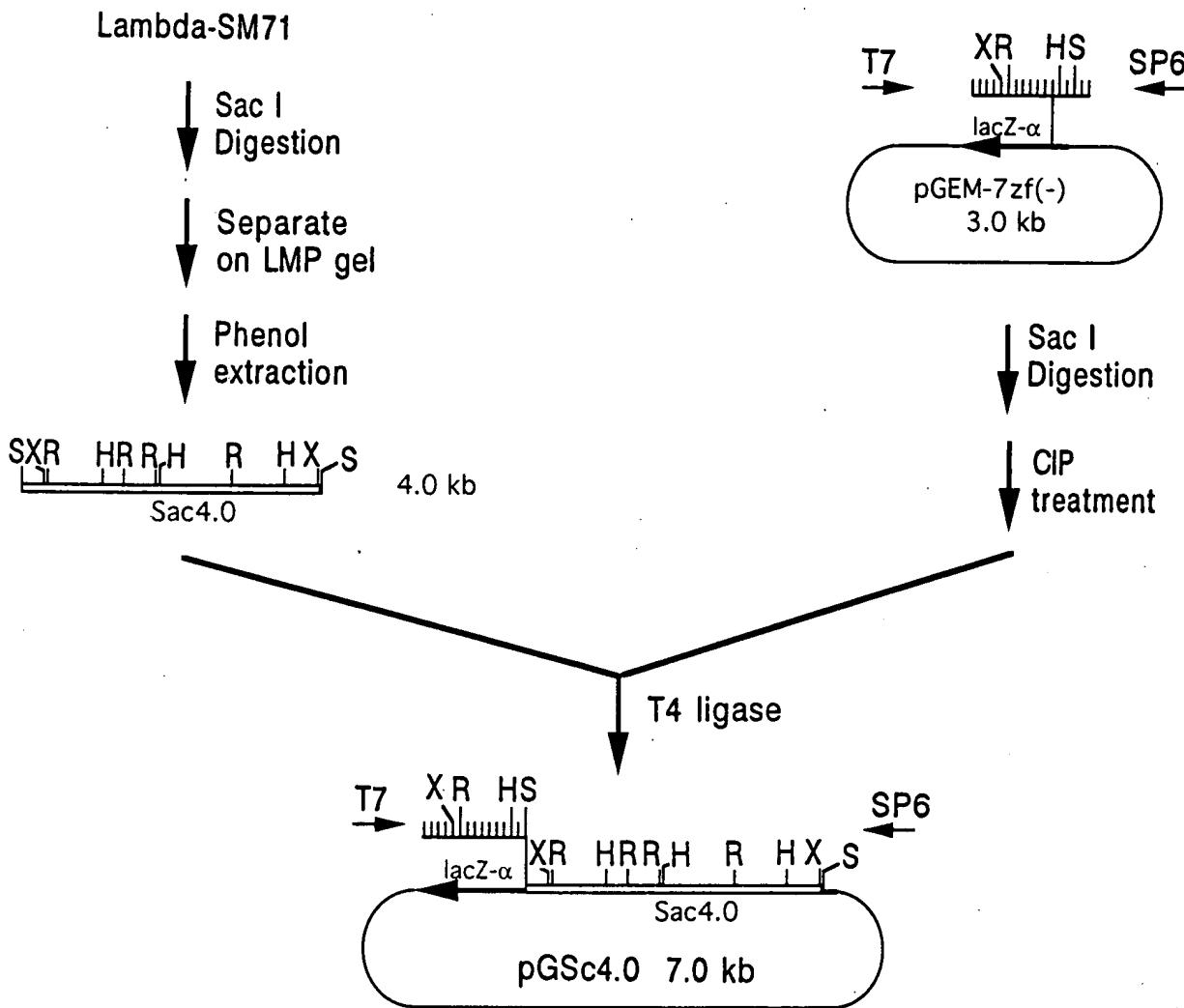
Map of Human *BCL-6* Locus





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**FIGURE 7**



Lambda-SM71 = a recombinant Lambda phage clone  
 containing *Bcl-6* breakpoint

H = Hind III

R = EcoR I

S = Sac I

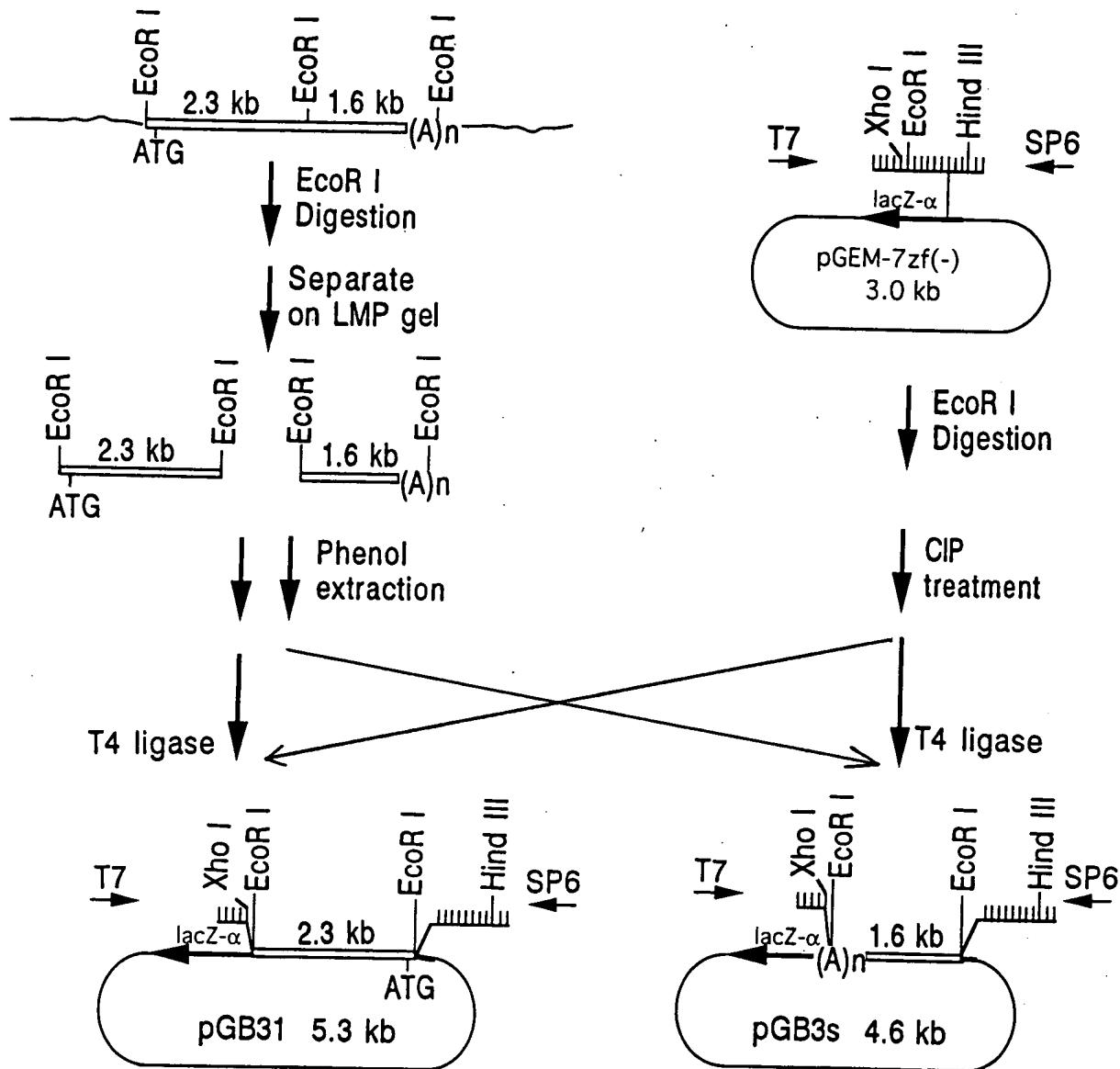
X = Xho I



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**FIGURE 8**

Lambda-B31





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FIGURE 9A

SEQ. ID. NOS. 1-2

## cdDNA and Amino Acid Sequences of BCL-6



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FIGURE 9B

A P G C E S R A F A P S L Y S T P P A S Y S M Y S H L P V S S L L F S D E 211

EFRDVRMpvANPFPKERALPCTDSARFVPTLEVSPP 1080  
EFRDVRMpvANPFPKERALPCTDSARFVPTLEVSPP 251

ANTIGENIC DETERMINANTIC ACID SEQUENCE OF THE HUMAN HEPATITIS C VIRUS 1200  
N V C H S N I Y S P K E T I P E E A R S D M H Y S V A E G L K P A A P S A R N A 291



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FIGURE 9C

TA G P T F A E E M G E T Q S E Y S D S S C E N G A F F C N E C D C R F S E A S 31



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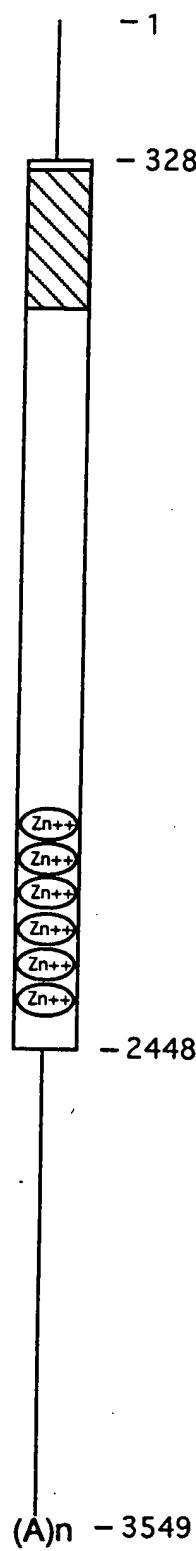
FIGURE 9D

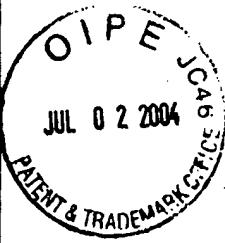


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### FIGURE 10A





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**FIGURE 10B**

1 MASPADSCIQ FTRHASDVLL NLNRLRSRDI LTDVVIVVSR EQFRAHKTVL  
51 MACSGLFYSI FTDQLKCNLS VINLDPEINP EGFCILLDFM YTSRLNLREG  
101 NIMAVMATAM YLQMEHVVDT CRKFIKASEA EMVSAIKPPR EEFLNSRMLM  
151 PQDIMAYRGR EVVENNLPLR SAPGCESRAF APSLYSGLST PPASYSMYSH  
201 LPVSSLLFSD EEFRDVRMPV ANPFPKERAL PCDSARPVPG EYSRPTLEVS  
251 PNVCHSNIYS PKETIPEEAR SDMHYSVAEG LKPAAPSARN APYFPCDKAS  
301 KEEERPSSED EIALHFEPPN APLNRKGLVS PQSPQKSDCQ PNSPTEACSS  
351 KNACILQASG SPPAKSPTDP KACNWKKYKF IVLNSLNQNA KPGGPEQAEI  
401 GRLSPRAYTA PPACQPPMEP ENLDLQSPTK LSASGEDSTI PQASRLNNIV  
451 NRSMTGSPRS SSESHSPLYM HPPKCTSCGS QSPQHAEMCL HTAGPTFAEE  
501 MGETQSEYSD SSCENGAFF [REDACTED] NECDCRESEE ASIKRHIIQI [REDACTED] SDKPYK [REDACTED]  
551 [REDACTED] QASAFRYKGN LASHKTVHTG EKPYR [REDACTED] NCG AOFNRPANLK THTRH [REDACTED] SGEK  
601 PYK [REDACTED] EYQVAHLRAH V1381TGEKPY PGE [REDACTED] GTRR [REDACTED] HLOTLKSHR  
651 [REDACTED] TGEKPYH [REDACTED] EKCNTHERHK SQRLHROK HGAITNTKVQ YRVSATDLPP  
701 ELPKAC\*



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FIGURE 11

ZFPJS	(2-56)	D G S F V	H V R Q L	Q E E N K O R E K G Q Y C O A	D V G G L V	F A H W A	A C C S H	F Q
KUP	(1-54)	D T A S H	H V L V Q	Q Q E F G F L C D G	A G D V Y	F A H W A	A C C S H	F Q
VASSR	(1-51)	M	N	Q E E N K O R E K G Q Y C O A	D V G G L V	F A H W A	A C C S H	F Q
ftk	(9-83)	C A R W N	H Q O N L	Q Q E F G F L C D G	A G D V Y	F A H W A	A C C S H	F Q
hatch	(182-186)	Q Y S N E	H R S F D A	Q Q E F G F L C D G	A G D V Y	F A H W A	A C C S H	F Q
PLZF	(10-63)	Q Q N P S	H G I C K A R	Q Q E F G F L C D G	A G D V Y	F A H W A	A C C S H	F Q
BCL-6	(8-62)	C Q F T R	H D L N H A	Q Q E F G F L C D G	A G D V Y	F A H W A	A C C S H	F Q
ZFPJS	(57-104)	D G	H Q	G G S V L P G F	D G S V L P G F	H Q	H Q	H Q
KUP	(55-107)	D	D	S E C I K Q P D	S E C I K Q P D	N L H	N L H	N L H
VASSR	(52-106)	N N	F I D S	N E Y E N L H	N E Y E N L H	E K H P I	E K H P I	E K H P I
ftk	(64-116)	S H	E E E R	S F E E R Q A R I T Q	S F E E R Q A R I T Q	L K O	L K O	L K O
hatch	(187-240)	S F	E E E R	H R N S Q H Y T L D F	H R N S Q H Y T L D F	R E L L C	R E L L C	R E L L C
PLZF	(64-114)	D Q L	K C N L S V I N L D	D Q L	D Q L	S K F Q Q L	S K F Q Q L	S K F Q Q L
BCL-6	(63-117)	N L D	D Q L	D Q L	D Q L	E E E R	E E E R	E E E R

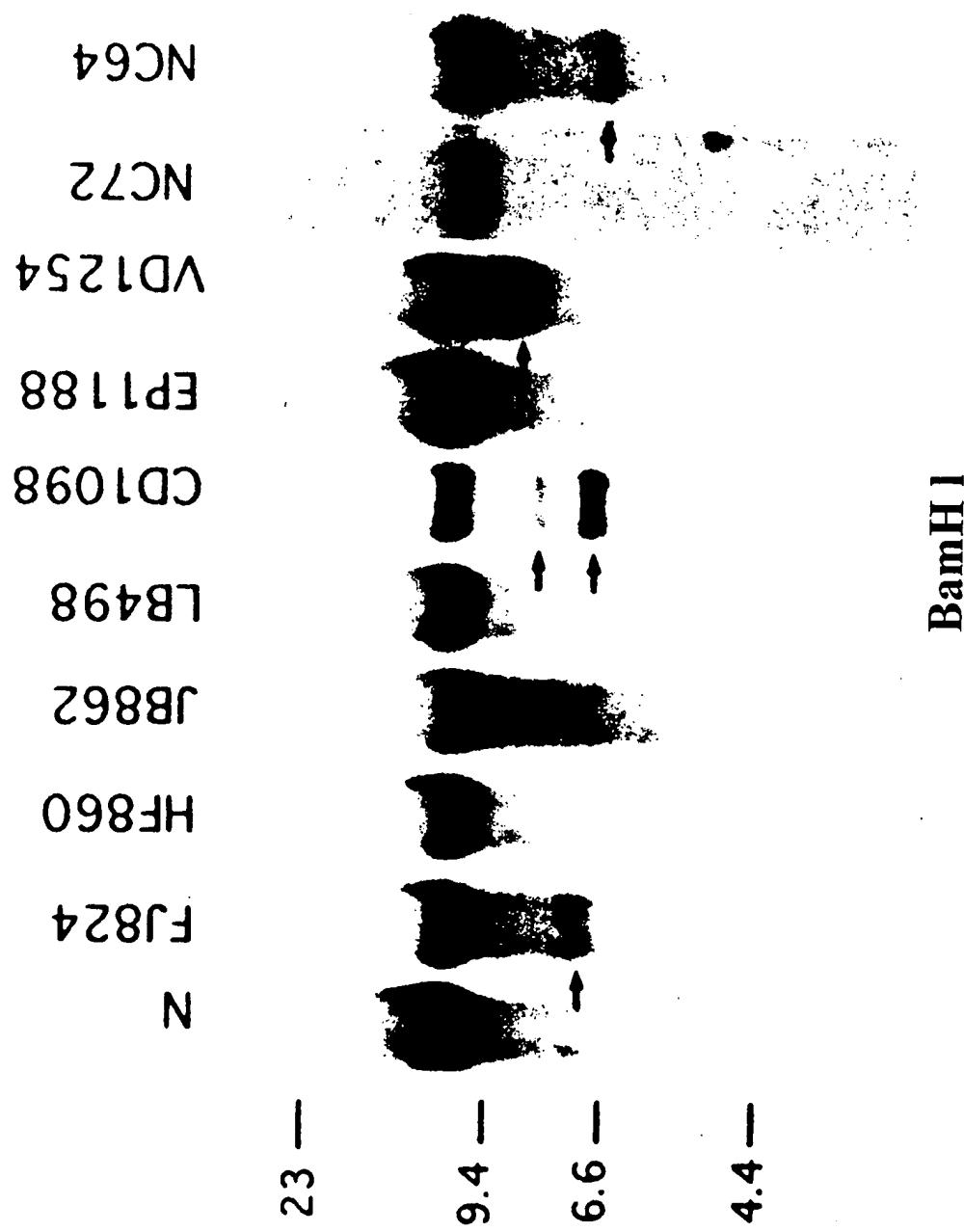


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FIGURE 13A



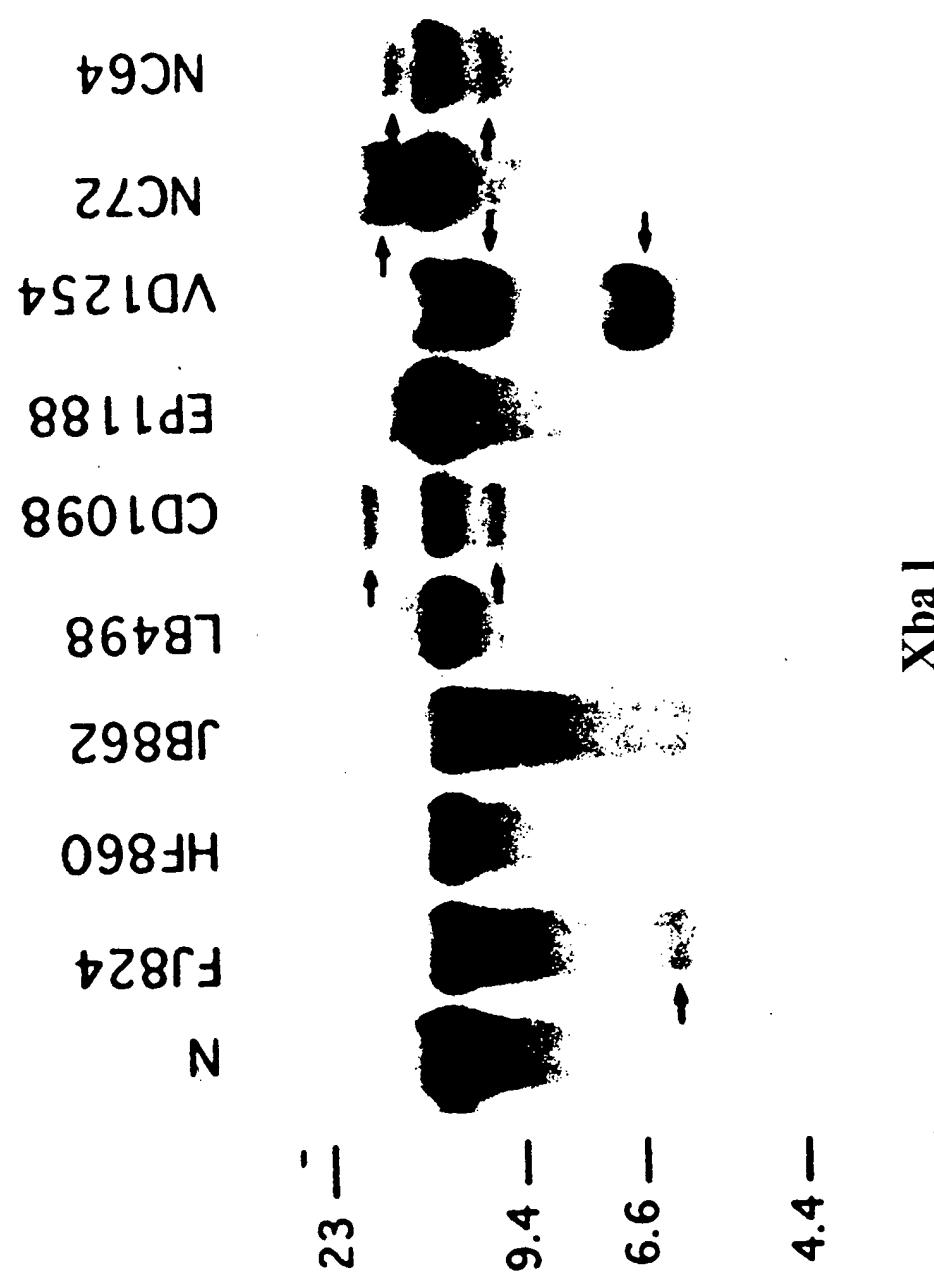


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FIGURE 13B





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FIGURE 14A

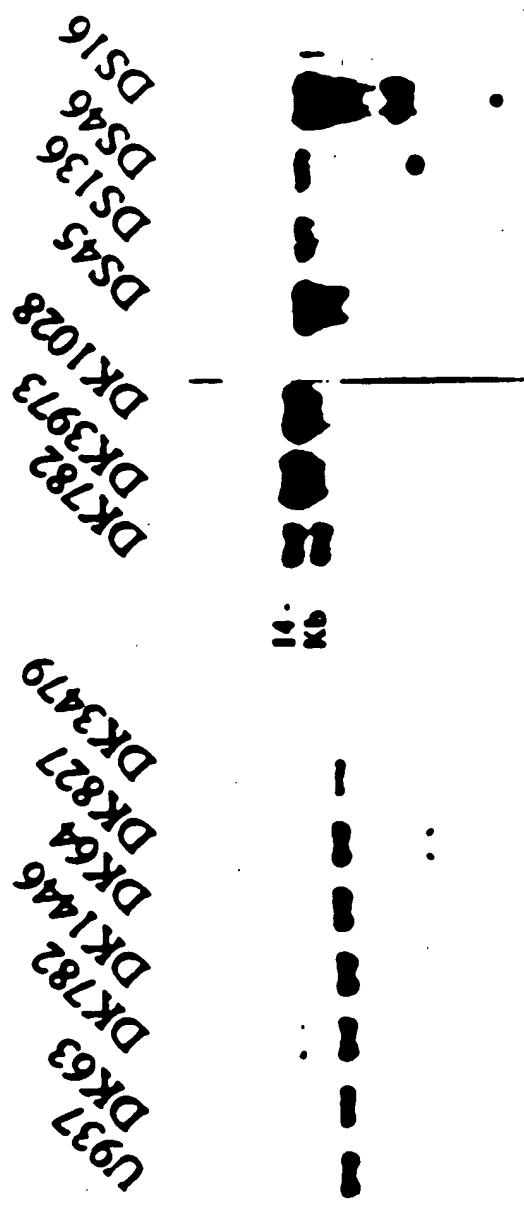


FIGURE 14B

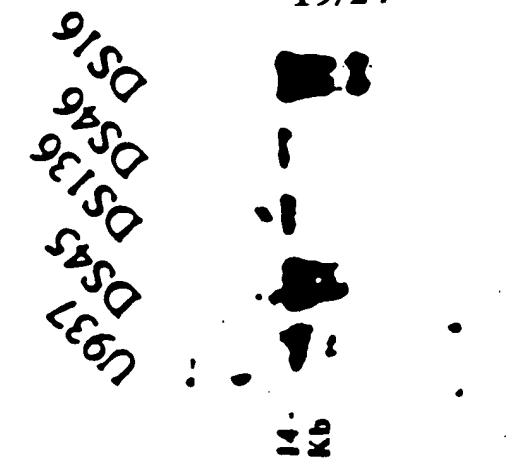
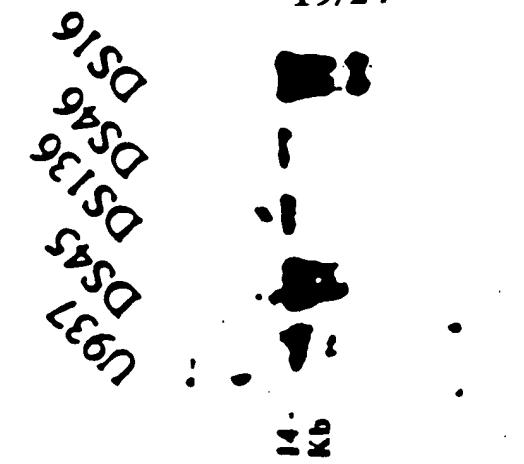


FIGURE 14C



*Bam*HII  
Sac4.0  
be

*Xba*I  
Sac4.0

*Xba*I  
Sac0.8





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FIGURE 16A

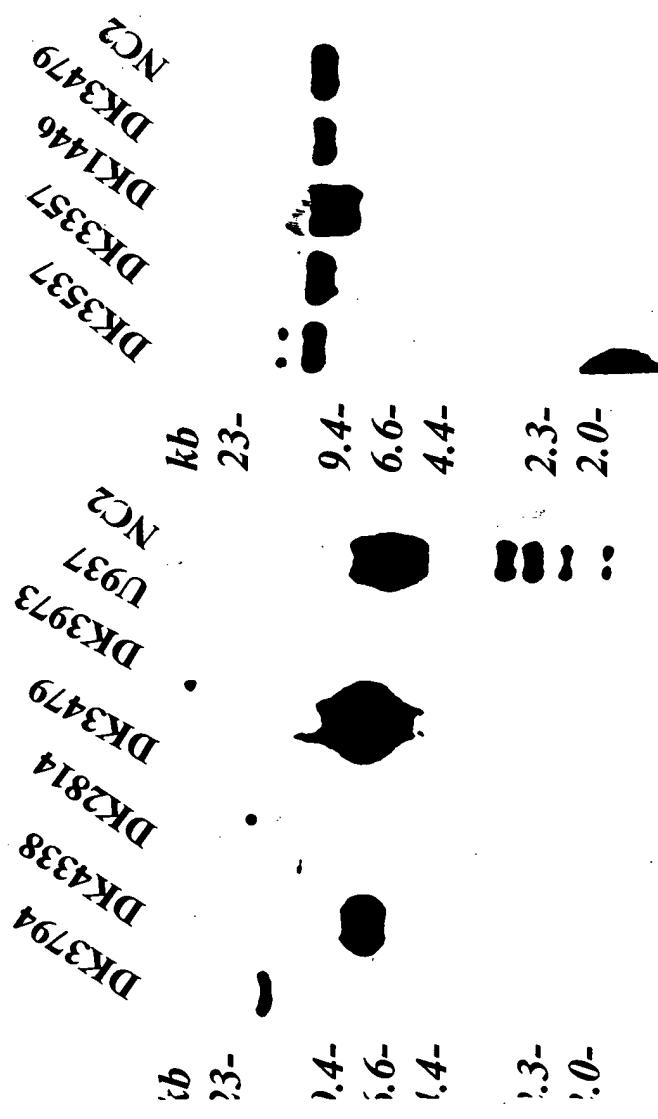
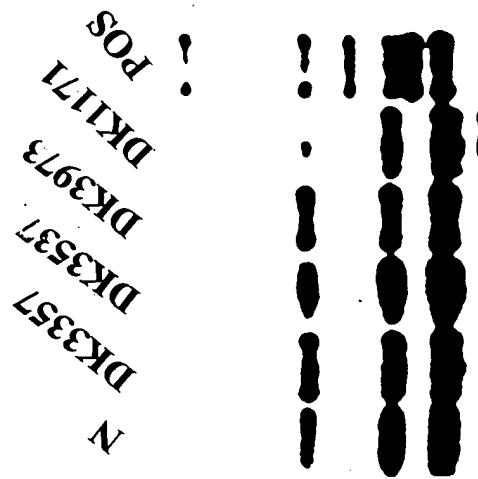


FIGURE 16C

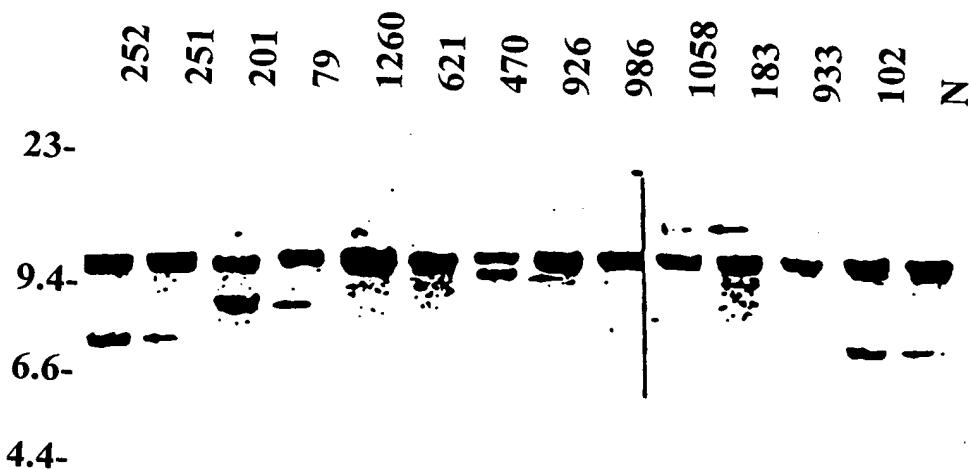


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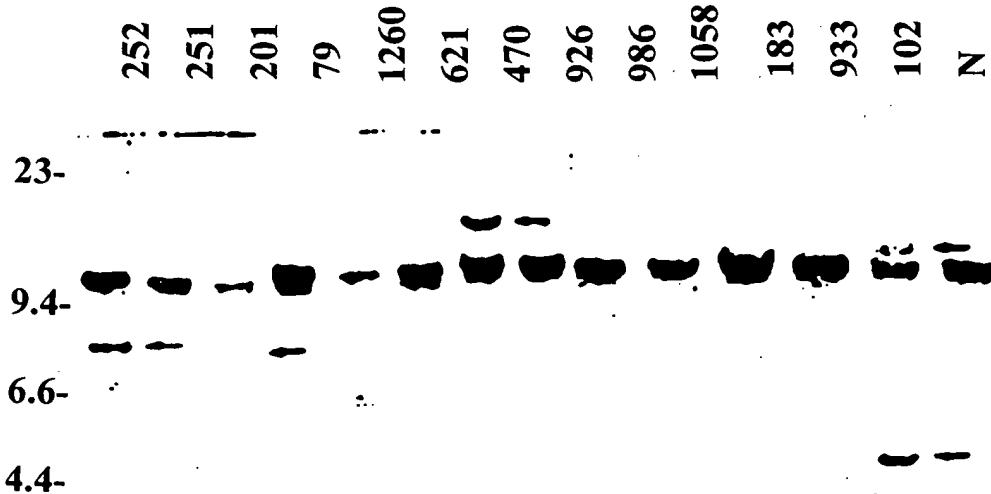
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**FIGURE 17A**



**FIGURE 17B**

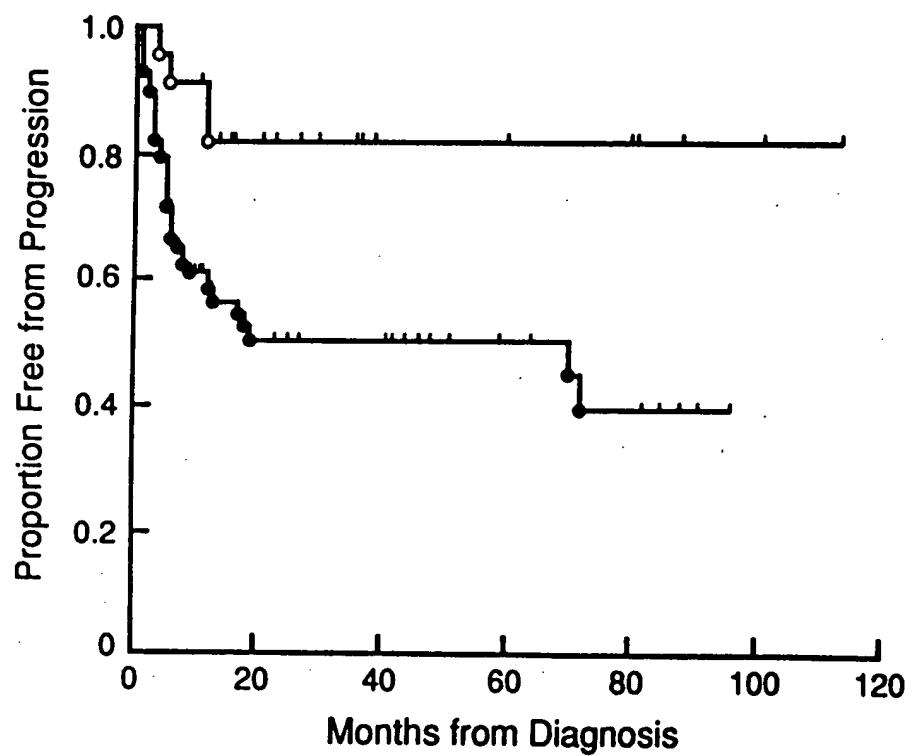




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**FIGURE 18A**





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**FIGURE 18B**

